3D1 heavy chain variable region sequence

٠ ټو٠	ATG M	GGT G	TGG W	AAC N	TGT C	ATC I	ATC I	TTC F	TTT F	30 CTG L	GTT V	ACA T	ACA T	GCT A	ACA T	GGT G	GTG V	CAC H	TCC S	60 CAG <u>Q</u>
	GTC V	CAG Q	CTG L	CAG Q	CAG Q	TCT S	GGG G	CCT P	GAG E	90 CTG L	GTG V	AGG R	CCT P	GGG G	GAA E	TCA S	GTG V	AAG K	ATT I	120 TCC S
	TGC C	aag K	GGT G	TCC S	GGC G	TAC Y	ACA T	TTC F	ACT T	150 GAT D	TAT Y	GCT A	ATA I	CAG O	, TGG W	GTG V	AAG K	CAG Q	AGT S	180 CAT H
-	GCA · A		AGT S		GAG E	TGG W	ATT I	GGA G	GTT V	210 ATT I	AAT N	ATT I	TAC Y	TAT Y	GAT D	AAT N	ACA T	AAC N	TAC Y	240 AAC <u>N</u>
D	CAG O	AAG K	ТТТ <u></u> F	AAG K	GGC G	AAG K	GCC A	ACA T	ATG M	270 ACT T	GTA V	GAC D	AAA K	TCC S	TCC S	AGC S	ACA T	GCC A	TAT Y	300 ATG M
	GAA E	CTT L	GCC A	AGA R	TTG L	ACA T	TCT S	GAG E	GAT D	330 TCT S	GCC A	ATC I	TAT Y	TAC Y	TGT C	GCA A	AGA R	GCG A	GCC A	360 TGG W
<u>=</u>	TAT <u>Y</u>	ATG M	GAC D	TAC Y	TGG W	GGT G	CAA Q	GGA G	ACC T	390 TCA S	GTC V	ACC T	GTC V	TCC S	TCA S		•			
						ï														

3D1 light chain variable region sequence

ATG	GAT	TCA	CAG	GCC	CAG	GTT	Conc	1. 2002	30										60
M	D	S	. CAG	A .	. CAG	v GTT	CTI L	' ATA I	TTG L	CTG L									
••	-	J	¥	**	~	٧		-	n	יי	. L	L	W	V	S	G	T	С	G
	٠								90										120
GAC	ATT	GTG	CTG	TCA	CAG	TCT	CCA	TCC	TCC	CTG	GCT	GTG	TCA	GCA	GGA	GAG	AAG	GTC	ACT
₽	I	V	L	S	Q	S	P	S	S	L	A	V	S	Α	G	E	K	V	T
ATG	AGC	TGC	AAA	TCC	AGT	CAC	» Cm	CITIC	150										180
M	S	C	K	S	S	O	AGT	L	CTC L										GCT
14	3	C	<u> </u>					<u></u> _	<u></u>	N	S	R	T	R	E	N	<u>Y</u>	<u>L</u> _	_ <u>A</u>
					. •				210										240
- TGG	TAC	CAG	CAG	AAA	CCA	GGG	CAG	TCT	CCT	AAA	CTG	CTG	ATC	TAC	TGG	GCA	TCC	ΔСТ	AGG
_W	Y.	Q	Q	K	P	G	Q	s	P	K	L	L	I	Y	W	A	s	ψ	R
									270										300
GAA	TCT	GGG	GTC	CCT	GAT	CGC	TTC	ACA	GGC	AGT	GGA	TCT	GGG	ACA	GAT	TTC	ACT	CTC	ACC
_E	<u>s</u>	G	V	P	D	R	F	${f T}$	G	S	G	· \$	G	T	D	F	${f T}$	L	T
•																			
ATC	AGC	AGT	GTG	CAG	GCT	GAA	C 2 C	000	330										360
I	S	S	V	Q	A	GAA E	GAC D	CTG		GTT								AAT	CTT
_	ے	3	V	Q	A	£	ע	L	A	V	Y	Y	С	<u>T</u>	0	S	<u>Y</u>	N	<u>L</u>
				٠.					390										
TAC	ACG	TTC	GGA	GGG	GGG	ACC .	AAG	CTG		ATA	AAA								
Y	\mathbf{T}	F	G	G	G	T	K	L	E	I	K		•		•				

Figure 1 (B)

Hu3D1 heavy chain variable region sequence

									30										60
AT	G GGI	TGG	AAC	TGI	ATC	ATC	TTC	TTT	CTG	GTT	ACC	: ACA	GCT	ACA	GGT	GTG	CAC	TCC	
М	G	W	N	С	I	I	F	F	L	V	т	T	A	. T	G	v	Н	s	
									_	•	-	•		· •	J	•			Ω
									90				-						120
GT	C CAG	CTG	GTG	CAG	TCT	GGG	GCT	GAG		AAG	AAG	י כריז	GGG	AGC	ጥር አ	GTG	אממ	GTG	
V	Q	L	v	Q	S	G	A	E	V	K	K	P	G	S	S	V	K	V	s
•	-	_	•	-	_	_	••	_	•	••		•	G	ی	٦	V	7	V	3
									150										180
TG	AAA	GCT	TCC	GGC	TAC	ACA	TTC	ACT		ጥልጥ	ርረጥ	מית	CAG	ጥርር	CTC	A C A	CAC	GCT	CCT
С	ĸ	A	s	G	Y	T	F	T	D	V	A	Т	0	W	V	R	Q	A	P
						_	_	-				-		••		K	Q	-	-
									210										240
- GGZ	A CAG	GGC	CTC	GAG	TGG	ATT	GGA	GTT		ידעע	<u>ን</u> ጥጥ	ጥልር	ጥልጥ	ርልጥ	מי מי מי	א כי א	ח ח כי	TAC	AAC
G	Q		L	E	W	I	G	v	I	N	T	Y	y	D	N	π	N	Y	N
•	~.	_	_			_	_					- -	-	<u> </u>	IV			<u> </u>	10
									270										300
⊒ =CA0	AAG	TTT	AAG	GGC	AAG	GCC	ACA	ATG		GΨΔ	GAC	מממ	TOC	ACG	AGC	ACA	GCC	TAT	ATG
<u></u> 0	к	F	ĸ	G	K	A	T	М	T	V	D	K	S	T	S	T	A	Y	M
Л -							-		_	V	ט	10	3	1	3	1	A	Ŧ	P4
U									330										360
JI GA.ª	CTT	AGT	TCT	TTG	AGA	ጥርጥ	GAG	GAT		CCC	CTT	ψ» m	ma c	TGT	CCN	202	000	666	
E	L	s	s	L	R	S	E	D	T	A	V	Y	Y	C		AGA			TGG
<u>,</u>		_		~	**			ט	1	A	V	1	1	C	A	R	<u>A</u>	_A	<u>W</u>
÷									390										
ייי בייי	ATG	GAC	ጥልሮ	тсс	GGT	C A A	CCT			GTC	1 CC	cmc		ma.					
= 'Y	M	D	Y	W	G	Q	G	T	L			GTC		TCA					
넵 ㅡ	11			**	G	Q	G	7	بد	V	Т	V	S	S					
U																			
Ą																	•		
_																			

Hu3D1 light chain variable region sequence

							•			3 ()									60
						C CAG		CTI	ATA	TTC	CTC	CTC	CT	A TGC	GTA	TCI	r GGC	ACC	TGI	oo OOO 1
	M	D	S	Q	A	Q	V	L	I	L	L	L	L	W	v	s	G	Ţ	C	. 666 G
			•															_		_
	GAC	א תאמו	CMC	· (m/					- 1	90)									120
	D D	I	V.	: CIC L	ACA T	A CAG	TCT	, CCY	. GAT	TCC	CTG	GCI	GTA	AGC	TTA	GGA	GAG	AGG	GCC	ACT
	==	-	V	ני	1	Q	S	P	D	S	Ŀ	A	V	S	L	G	E	R	A	${f T}$
										150								•	•	
	ATT	AGC	TGC	AAA	TCC	AGT	CAG	AGT	СТС	CTC	ם מ מ	λCm	A C A	N.C.C	003	~ ~				180
	I	s	С	<u>K</u>	s_	S	0	_ s	L	L	N	S	R	ACC T	R	GAG E			_	_
	·															<u> </u>	N	<u>Y</u>	L_	<u>A</u>
										210										240
	TGG	TAC	CAG	CAG	AAA	CCA	GGG	CAG	CCT	CCT	AAA	CTG	CTG	ATC	TAC	TGG	GCA	TCC	Δ (Τ	2 4 0
	_ W	Y	Q	. Q	K	P	G	Q	P .	· P	K	L	L	I	Y	W	A	s	T	. R
		•								٠										
	ממם	ጥርብ	GGG	CTTC	CCM	C N M	000			270										300
_	E	_S	G	V	P	GAT D	R	TTC								GAT	TTC	ACT	CTC	ACC
	-	_=			-	ע	K	r	S	G	S	G	S	G	${f T}$	D	F	T	L	${f T}$
ب ج	i					•				330										•
	ATC .	AGC	AGT	CTG	CAG	GCT	GAA	GAC			Cum	TAT	TAC	mc c	N C C	~~~				360
≓	I	S	s	L	Q	A	E	D	V	A	V	Y	Y	C .	ACG '	CAA O			AAT	
											•	-	_	C	<u> </u>	<u> </u>	S	<u> </u>	_N	<u>L</u>
										390										•
	TAC A				CAG	GGG .	ACC .	AAG (GTG (SAA .	ATA A	AAA								
	<u>Y</u>	T	F	G	Q	G	${f T}$	K	V	E	I	K								
٠.																				

Competition Binding Assay of Anti-B7.2 mAbs

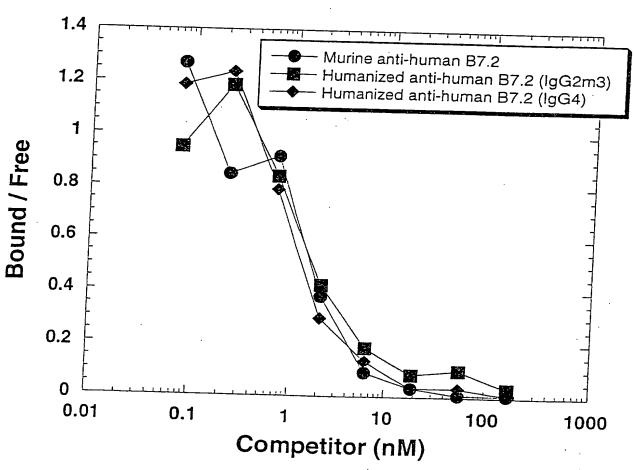


Figure 3

Direct Binding Assay of Anti-B7.2 mAbs

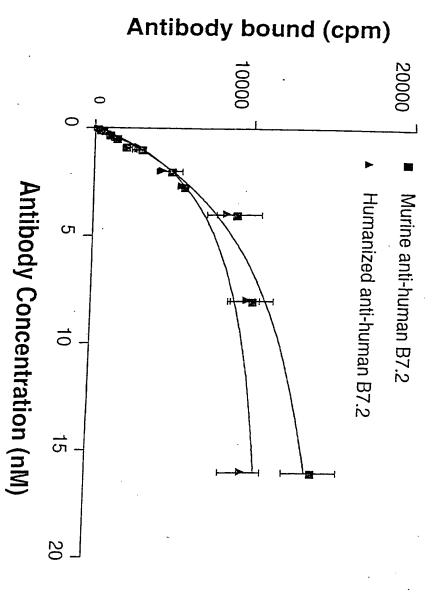


Figure 4

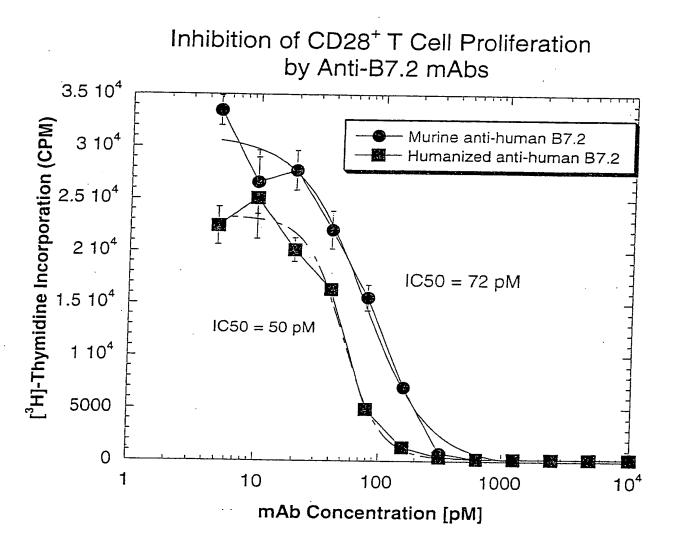


Figure 5

Inhibition of a Mixed Lymphocyte Reaction by Anti-B7 Antibodies and CTLA4IG

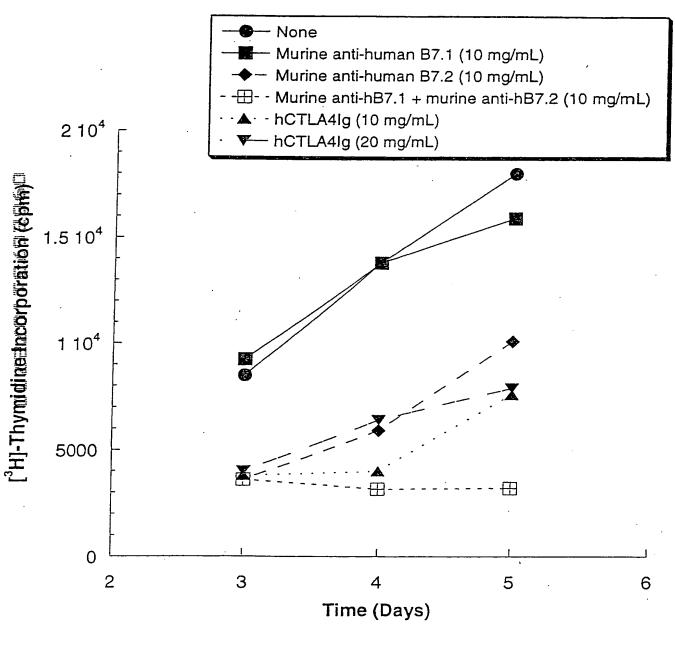


Figure 6

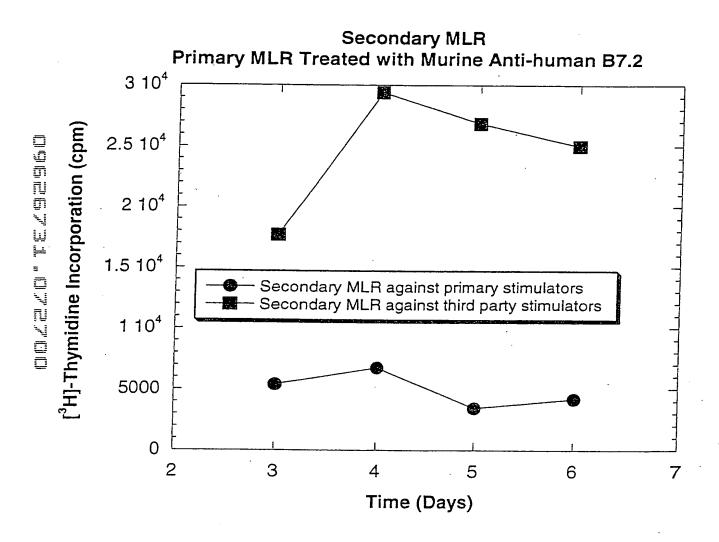


Figure 7

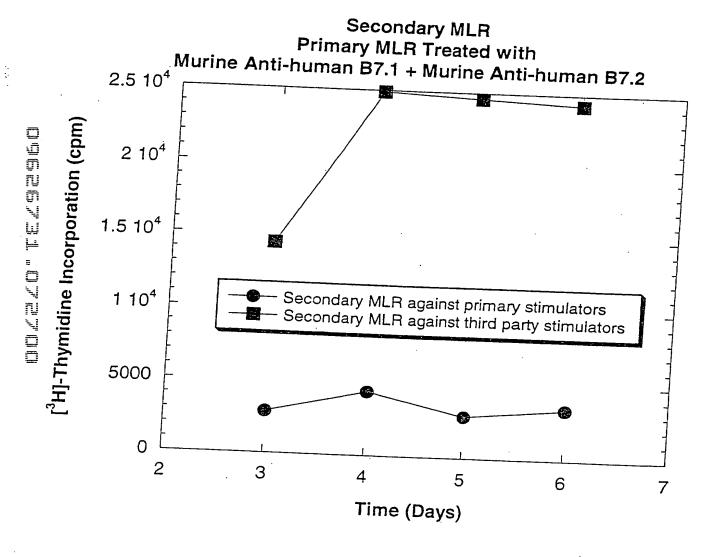
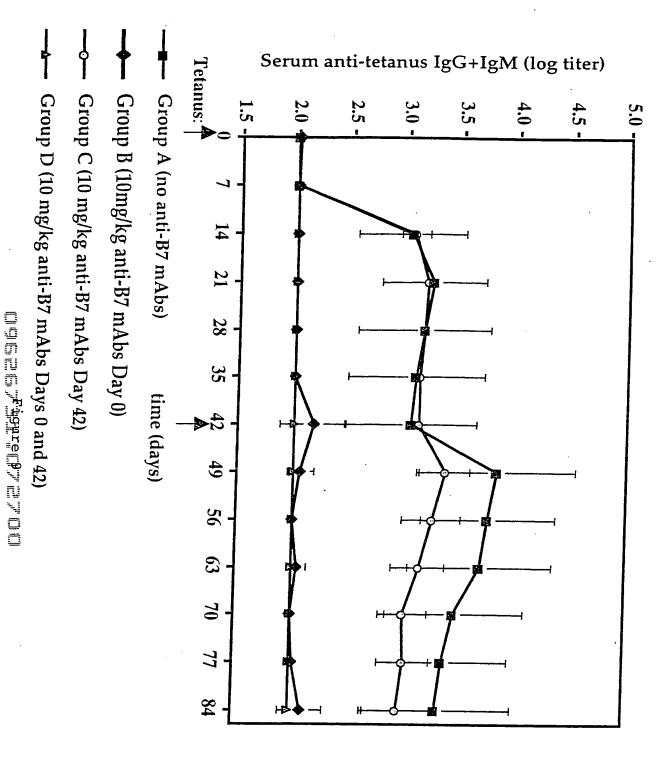


Figure 8

Antibody Response to Tetanus Immunization During Costimulation Blockade with Humanized Anti-B7.1 and Anti-B7.2



Anti-B7-2, Serum concentration



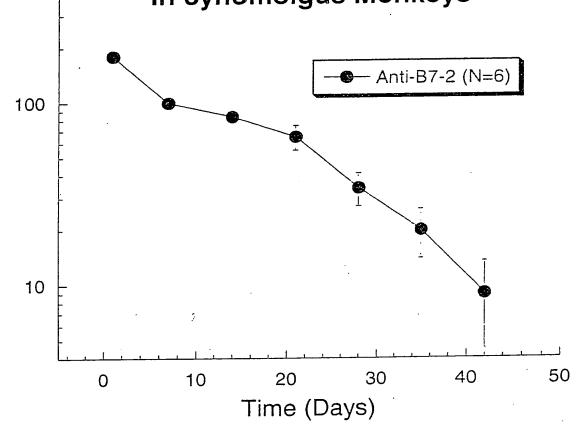


Figure 10